1/7

GATGGGCCTCANGTTCAT-3' 5'-ATGAACCGGAGGCCCATC-3' (%) (8) (64 (64 (22)26 69 (83) 55 5 O (8) 54 ග (L) (e1 49 (23)Ø <u>.</u> 80 (32) TARGET DNA SEQUENCE NO.65 8 **24** (2) (34) 9 O 8 (K) (8)7 വ တ × GATGGGGCTCGNGTTCAT GATGGGGCTCCNGTTCAT 5'-GATGGGACTCANGTTCAT GATGGGGCTCANGTTCAT GATGGGGCTCTNGTTCAT GATGGGACTCGNGTTCAT GATGGGACTCCNGTTCAT GATGGGACTC<u>IN</u>GTTCAT FIG.

GATGGGCCTCGNGTTCAT GATGGGICTCGNGTTCAT GATGGG<u>C</u>CTC<u>CN</u>GTTCAT GATGGGCCTCINGTTCAT GATGGGICTCANGTTCAT GATGGGICTCCNGTTCAT GATGGGICTCINGTTCAT



FULL MATCH



TWO-BASE MISMATCH



THREE-BASE MISMATCH